04C0
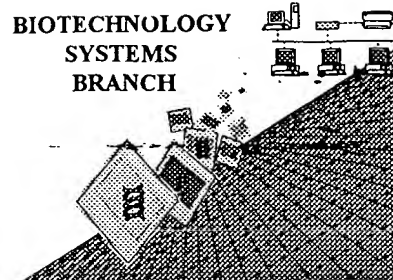
RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _09/641,931_

Source: _OIPE_

Date Processed by STIC: _8/29/2000_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**

2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

 FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENCER, 703-308-4212.

TO **REDUCE** ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October-1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).
Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
http://www.uspto.gov/web/offices/pac/checker

OIPE

pp 1-5

RAW SEQUENCE LISTING   DATE: 08/29/2000
PATENT APPLICATION: US/09/641,931   TIME: 12:54:49

Input Set : A:\ES.txt
Output Set: N:\CRF3\08292000\I641931.raw

Does Not Comply
Corrected Diskette Needed

```
       3 <110> APPLICANT: PHENOGENE THERAPEUTIQUES INC.
       5 <120> TITLE OF INVENTION: NUCLEIC ACID MOLECULE, METHOD AND KIT FOR SELECTING A NUCLEIC ACID HAVING
       6       A DESIRED FEATURE
       8 <130> FILE REFERENCE: 29313-001
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/641,931
C--> 10 <141> CURRENT FILING DATE: 2000-08-18
      10 <160> NUMBER OF SEQ ID NOS: 32
      12 <170> SOFTWARE: PatentIn version 3.0
      14 <210> SEQ ID NO: 1
      15 <211> LENGTH: 24
      16 <212> TYPE: DNA
      17 <213> ORGANISM: artificial sequence
W--> 19 <220> FEATURE:
W--> 19 <223> OTHER INFORMATION:
      19 <400> SEQUENCE: 1
      20 ggatccaata gaggattctt taac                          24
      23 <210> SEQ ID NO: 2
      24 <211> LENGTH: 21
      25 <212> TYPE: DNA
      26 <213> ORGANISM: artificial sequence
W--> 28 <220> FEATURE:
W--> 28 <223> OTHER INFORMATION:
      28 <400> SEQUENCE: 2
      29 tcaccactct tctgtccctt c                             21
      32 <210> SEQ ID NO: 3
      33 <211> LENGTH: 25
      34 <212> TYPE: DNA
      35 <213> ORGANISM: artificial sequence
W--> 37 <220> FEATURE:
W--> 37 <223> OTHER INFORMATION:
      37 <400> SEQUENCE: 3
      38 ggatcctacg aacatgcgac cactg                         25
      41 <210> SEQ ID NO: 4
      42 <211> LENGTH: 21
      43 <212> TYPE: DNA
      44 <213> ORGANISM: artificial sequence
W--> 46 <220> FEATURE:
W--> 46 <223> OTHER INFORMATION:
      46 <400> SEQUENCE: 4
      47 tcatcttcgt gtgctagtca g                             21
      50 <210> SEQ ID NO: 5
      51 <211> LENGTH: 30
      52 <212> TYPE: DNA
      53 <213> ORGANISM: artificial sequence
W--> 55 <220> FEATURE:
W--> 55 <223> OTHER INFORMATION:
      55 <400> SEQUENCE: 5
```

(global error)

insert there mandatory numeric identifiers wherever <213> response is Unknown or Artificial Sequence

```
          RAW SEQUENCE LISTING              DATE: 08/29/2000
          PATENT APPLICATION:  US/09/641,931   TIME: 12:54:49

               Input Set : A:\ES.txt
               Output Set: N:\CRF3\08292000\I641931.raw

       56 agcgaattcg tcctgtggac agatcactgc                         30
       59 <210> SEQ ID NO: 6
       60 <211> LENGTH: 30
       61 <212> TYPE: DNA
       62 <213> ORGANISM: artificial sequence
W--> 64 <220> FEATURE:
W--> 64 <223> OTHER INFORMATION:
       64 <400> SEQUENCE: 6
       65 gctctcgagg aaggcacagc tgctttccac                         30
       68 <210> SEQ ID NO: 7
       69 <211> LENGTH: 30
       70 <212> TYPE: DNA
       71 <213> ORGANISM: artificial sequence
W--> 73 <220> FEATURE:
W--> 73 <223> OTHER INFORMATION:
       73 <400> SEQUENCE: 7
       74 cttctcgagc agtttaaacg tgagcttccc                         30
       77 <210> SEQ ID NO: 8
       78 <211> LENGTH: 30
       79 <212> TYPE: DNA
       80 <213> ORGANISM: artificial sequence
W--> 82 <220> FEATURE:
W--> 82 <223> OTHER INFORMATION:
       82 <400> SEQUENCE: 8
       83 acgtctagat catcttcgtg tgctagtcag                         30
       86 <210> SEQ ID NO: 9
       87 <211> LENGTH: 47
       88 <212> TYPE: DNA
       89 <213> ORGANISM: artificial sequence
W--> 91 <220> FEATURE:
W--> 91 <223> OTHER INFORMATION:
       91 <400> SEQUENCE: 9
       92 tcgagcagat ctgcagcacc actggtcacg gcaatgtgtc ggagcgg      47
       95 <210> SEQ ID NO: 10
       96 <211> LENGTH: 43
       97 <212> TYPE: DNA
       98 <213> ORGANISM: artificial sequence
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W--> 100 <223> OTHER INFORMATION:
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       101 ccgctccgac acattgccgt gaccagtggt gctgcagatc tgc          43
       104 <210> SEQ ID NO: 11
       105 <211> LENGTH: 63
       106 <212> TYPE: DNA
       107 <213> ORGANISM: artificial sequence
W--> 109 <220> FEATURE:
W--> 109 <223> OTHER INFORMATION:
       109 <400> SEQUENCE: 11
       110 gtgtccaagc catcagaggg gaaataaagc atctctacgg tggtcctaaa tagtcagcat  60
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       112 agt                                                    63
       115 <210> SEQ ID NO: 12
       116 <211> LENGTH: 28
       117 <212> TYPE: DNA
       118 <213> ORGANISM: artificial sequence
W--> 120 <220> FEATURE:
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       120 <400> SEQUENCE: 12
       121 ccagagctca tgcggaccac tcttctgt                         28
       124 <210> SEQ ID NO: 13
       125 <211> LENGTH: 24
       126 <212> TYPE: DNA
       127 <213> ORGANISM: artificial sequence
W--> 129 <220> FEATURE:
W--> 129 <223> OTHER INFORMATION:
       129 <400> SEQUENCE: 13
       130 tcgcgattta aattaattaa gctt                             24
       133 <210> SEQ ID NO: 14
       134 <211> LENGTH: 24
       135 <212> TYPE: DNA
       136 <213> ORGANISM: artificial sequence
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W--> 138 <223> OTHER INFORMATION:
       138 <400> SEQUENCE: 14
       139 aagcttaatt aatttaaatc gcga                             24
       142 <210> SEQ ID NO: 15
       143 <211> LENGTH: 18
       144 <212> TYPE: DNA
       145 <213> ORGANISM: artificial sequence
W--> 147 <220> FEATURE:
W--> 147 <223> OTHER INFORMATION:
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       148 agacgcgtag atctcacc                                   18
       151 <210> SEQ ID NO: 16
       152 <211> LENGTH: 20
       153 <212> TYPE: DNA
       154 <213> ORGANISM: artificial sequence
W--> 156 <220> FEATURE:
W--> 156 <223> OTHER INFORMATION:
       156 <400> SEQUENCE: 16
       157 gatccgcacc gcaatatggc                                 20
       160 <210> SEQ ID NO: 17
       161 <211> LENGTH: 25
       162 <212> TYPE: DNA
       163 <213> ORGANISM: artificial sequence
W--> 165 <220> FEATURE:
W--> 165 <223> OTHER INFORMATION:
       165 <400> SEQUENCE: 17
       166 tctagagatg cattatgcac atcag                            25
```

```
                    RAW SEQUENCE LISTING              DATE: 08/29/2000
                    PATENT APPLICATION:  US/09/641,931    TIME: 12:54:49

                    Input Set : A:\ES.txt
                    Output Set: N:\CRF3\08292000\I641931.raw

        169 <210> SEQ ID NO: 18
        170 <211> LENGTH: 60
        171 <212> TYPE: DNA
        172 <213> ORGANISM: artificial sequence
W--> 174 <220> FEATURE:
W--> 174 <223> OTHER INFORMATION:
        174 <400> SEQUENCE: 18
        175 tccaagccat cagaggggaa ataaagcatc tctacggtgg tcctaaatag tcagcatagt     60
        178 <210> SEQ ID NO: 19
        179 <211> LENGTH: 60
        180 <212> TYPE: DNA
        181 <213> ORGANISM: artificial sequence
W--> 183 <220> FEATURE:
W--> 183 <223> OTHER INFORMATION:
        183 <400> SEQUENCE: 19
        184 actatgctga ctatttagga ccaccgtaga gatgctttat ttccctctg atggcttgga      60
        187 <210> SEQ ID NO: 20
        188 <211> LENGTH: 20
        189 <212> TYPE: DNA
        190 <213> ORGANISM: artificial sequence
W--> 192 <220> FEATURE:
W--> 192 <223> OTHER INFORMATION:
        192 <400> SEQUENCE: 20
        193 tagtcagcat agtacatttc                                                 20
        196 <210> SEQ ID NO: 21
        197 <211> LENGTH: 51
        198 <212> TYPE: DNA
        199 <213> ORGANISM: artificial sequence
W--> 201 <220> FEATURE:
W--> 201 <223> OTHER INFORMATION:
        201 <400> SEQUENCE: 21
        202 tcgatccgaa ttcgcggccg ctctattgga tcctcgagca gatctgcagc a             51
        205 <210> SEQ ID NO: 22
        206 <211> LENGTH: 148
        207 <212> TYPE: DNA
        208 <213> ORGANISM: artificial sequence
W--> 210 <220> FEATURE:
W--> 210 <223> OTHER INFORMATION:
        210 <400> SEQUENCE: 22
        211 agatgaatca agcttatcga taccgtcgag catgcatcta ggtgtccaag ccatcagagg     60
        213 ggaaataaag catctctacg gtggtcctaa atagtcagca tagtacattt catctgacta     120
        215 atactacaac accaccacca tgaataga                                        148
        218 <210> SEQ ID NO: 23
        219 <211> LENGTH: 18
        220 <212> TYPE: DNA
        221 <213> ORGANISM: artificial sequence
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W--> 223 <223> OTHER INFORMATION:
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      224 gagtggtccg catggtga                                            18
      227 <210> SEQ ID NO: 24
      228 <211> LENGTH: 54
      229 <212> TYPE: DNA
      230 <213> ORGANISM: artificial sequence
W--> 232 <220> FEATURE:
W--> 232 <223> OTHER INFORMATION:
      232 <400> SEQUENCE: 24
      233 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aagggggaatt tcgcgattta aatt      54
      236 <210> SEQ ID NO: 25
      237 <211> LENGTH: 48
      238 <212> TYPE: DNA
      239 <213> ORGANISM: Sindbis virus
      241 <400> SEQUENCE: 25
      242 tctgcagcac cactggtcac ggcaatgtgt ttgctcggaa atgtgagc             48
      245 <210> SEQ ID NO: 26
      246 <211> LENGTH: 16
      247 <212> TYPE: PRT
      248 <213> ORGANISM: Sindbis virus
      250 <400> SEQUENCE: 26
      252 Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val Ser
      253 1               5                   10                  15
      255 <210> SEQ ID NO: 27
      256 <211> LENGTH: 48
      257 <212> TYPE: DNA
      258 <213> ORGANISM: artificial sequence
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W--> 260 <223> OTHER INFORMATION:
      260 <400> SEQUENCE: 27
      261 tctgcagcac cactggtcac ggcaatgtgt cggagcggaa atgtgagc             48
      264 <210> SEQ ID NO: 28
      265 <211> LENGTH: 16
      266 <212> TYPE: PRT
      267 <213> ORGANISM: artificial sequence
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W--> 269 <223> OTHER INFORMATION:
      269 <400> SEQUENCE: 28
      271 Ser Ala Ala Pro Leu Val Thr Ala Met Cys Arg Ser Gly Asn Val Ser
      272 1               5                   10                  15
      274 <210> SEQ ID NO: 29
      275 <211> LENGTH: 44
      276 <212> TYPE: DNA
      277 <213> ORGANISM: artificial sequence
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W--> 278 <223> OTHER INFORMATION:
      279 <400> SEQUENCE: 29
      280 gagagagaga gagtttaaac gtcgactttt tttttttttt tttt                 44
      283 <210> SEQ ID NO: 30
      284 <211> LENGTH: 34
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VERIFICATION SUMMARY   DATE: 08/29/2000
PATENT APPLICATION: US/09/641,931   TIME: 12:54:50

        Input Set : A:\ES.txt
        Output Set: N:\CRF3\08292000\I641931.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:19 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:19 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:28 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:28 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:37 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:37 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:46 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:46 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:64 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:73 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:82 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:82 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:138 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:147 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:210 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:223 M:258 W: Mandatory Feature missing, <220> FEATURE:
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VERIFICATION SUMMARY   DATE: 08/29/2000
PATENT APPLICATION: US/09/641,931   TIME: 12:54:50

Input Set : A:\ES.txt
Output Set: N:\CRF3\08292000\I641931.raw

L:232 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:232 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:260 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:260 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:

Raw Sequence Listing Error Summary

ERROR DETECTED   SUGGESTED   CORRECTION   SERIAL NUMBER: 09/641,931

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 _____ Wrapped Nucleics
The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".

2 _____ Wrapped Aminos
The amino acid number/text at the end of each line "wrapped " down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".

3 _____ Incorrect Line Length
The rules require that a line not exceed 72 characters in length. This includes spaces.

4 _____ Misaligned Amino Acid Numbering
The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs
between the numbering. It is recommended to delete any tabs and use spacing between the numbers.

5 _____ Non-ASCII
This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.

6 _____ Variable Length
Sequence(s) _____ contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and
indicate in the (ix) feature section that some may be missing.

7 _____ PatentIn ver. 2.0 "bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
sequence(s) _____. Normally, PatentIn would automatically generate this section from the
previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section
to the subsequent amino acid sequence. **This applies primarily to the mandatory <220>-<223>
sections for Artificial or Unknown sequences.**

8 _____ Skipped Sequences (OLD RULES)
Sequence(s) _____ missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).

9 _____ Skipped Sequences (NEW RULES)
Sequence(s) _____ missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

10 _____ Use of n's or Xaa's (NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

11 _____ Use of <213>Organism (NEW RULES)
Sequence(s) _____ are missing this mandatory field or its response.

1-24, 27-29 (maybe more)

12 _____ Use of <220>Feature (NEW RULES)
Sequence(s) _____ are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of new Rules)

13 _____ PatentIn ver. 2.0 "bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted
file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.